ENTERED

PCT

```
 1 <110> APPLICANT: Choi, Eui-Sung
 2        Rhee, Sang-Ki
 3        Sohn, Jung-Hoon
 4        Park, Soo-Dong
 5        Lee, Yoon-Hyoung
 6        Lee, Seung-Jae
 7        Jang, Jae-Kweon
 8        Choi, Seok-Keun
 9        Son, Young-Rok
10 <120> TITLE OF INVENTION: VECTOR FOR THE TRANSFORMATION OF PHAFFIA
11        RHODOZYMA AND PROCESS OF TRANSFORMATION THEREBY
12 <130> FILE REFERENCE: 118.12-US-WO
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/830,691A
14 <141> CURRENT FILING DATE: 2001-04-26
15 <150> PRIOR APPLICATION NUMBER: KR 1998/46547
16 <151> PRIOR FILING DATE: 1998-10-31
17 <150> PRIOR APPLICATION NUMBER: PCT/KR99/00265
18 <151> PRIOR FILING DATE: 1999-05-29
19 <160> NUMBER OF SEQ ID NOS: 20
20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 1223
24 <212> TYPE: DNA
25 <213> ORGANISM: Phaffia rhodozyma
26 <400> SEQUENCE: 1
27        atggtcaacg ttcccaagac tcgacgtgag ttatagcaat ttcaacaact ctccagacga 60
28        caaatattcc agtgcatcga aagagtttgt ggataaacgc gacagtttca agggaaagag 120
29        tcgatggaca gatttggaag acttagccgg tcaaggaact tggggatcac gtggcggagg 180
30        actcatcaga agaagtcggg atttgtttga tcatagtggg atcaagacaa actggaggat 240
31        atggctcgcc ttggaaggga atctccggcc tggattcgag gatccgaaag ttgtacgtat 300
32        ggaaaagctt acacggcttg gatttattat ctttcatagg aacctactgc aagggtaagg 360
33        cttgcaagaa gcacacgtaa gtcgcttatc ctctccactc tttcatggca tattgtcaac 420
34        gactggacaa cgcgtccgtt ttgaaacaag tgacttacct gtgaaatttg attctacacc 480
35        tgtatttagc cctcacaagg tacatatcac atcctcccac cccaccctgc ccaacttctt 540
36        cagttcatct tgctctcggt ttccacattc cctgatgacc tccttgtatg ttctttgcga 600
37        acgtttgttt ctgtttctgt aggtgaccca gtacaagaag ggaaaggact ccatcttcgc 660
38        ccagggaaag cgacgatacg accgaaagca gtccggttac ggaggtcaga ccaagcccgt 720
39        tttccacaag aaggctaaga ccaccaagaa ggtcgtcctt cgattggcgg tattttttgtt 780
40        tattttgaat tcttttttgtg tatgcagact tttgatgatt atgctcctct gtcgtttttt 840
41        ctcttcaaac agagtgctcc gtctgcagtt cgttcttcct tccaaccaaa acttcaacta 900
42        cagacatcat aaacagacat cttacttcgg tgttctctct tttttccgc agagtacaag 960
43        atgcagatga ccctcaagcg atgcaagcac ttcgagcttg gaggagacaa gaagaccaag 1020
44        ggttcgtctt ttgtccatat attctctggt tcacttctta tgttcctaac gtacttgttt 1080
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45         ccttttttggt tcggatgttg tttctatcgg tggtgttttc ttttctttgg atgcattatc 1140
46         atttatcgtg ttggactgtt ttcctctgct cgtttctttc tcctctgtac ttgtgcttct 1200
47         caggagccgc catctctttc taa                                        1223
49 <210> SEQ ID NO: 2
50 <211> LENGTH: 350
51 <212> TYPE: DNA
52 <213> ORGANISM: Phaffia rhodozyma
53 <220> FEATURE:
54 <221> NAME/KEY: CDS
55 <222> LOCATION: (30)...(347)
56 <400> SEQUENCE: 2
57         cccttcaagt ctcgtctcaa tcagtcaag atg gtc aac gtt ccc aag act cga     53
58                                          Met Val Asn Val Pro Lys Thr Arg
59                                           1               5
60         cga acc tac tgc aag ggt aag gct tgc aag aag cac acc cct cac aag    101
61         Arg Thr Tyr Cys Lys Gly Lys Ala Cys Lys Lys His Thr Pro His Lys
62              10              15                  20
63         gtg acc cag tac aag aag gga aag gac tcc atc ttc gcc cag gga aag    149
64         Val Thr Gln Tyr Lys Lys Gly Lys Asp Ser Ile Phe Ala Gln Gly Lys
65          25                  30                  35                  40
66         cga cga tac gac cga aag cag tcc ggt tac gga ggt cag acc aag ccc    197
67         Arg Arg Tyr Asp Arg Lys Gln Ser Gly Tyr Gly Gly Gln Thr Lys Pro
68                      45                  50                  55
69         gtt ttc cac aag aag gct aag acc acc aag aag gtc gtc ctt cga ttg    245
70         Val Phe His Lys Lys Ala Lys Thr Thr Lys Lys Val Val Leu Arg Leu
71                  60                  65                  70
72         gag tgc tcc gtc tgc aag tac aag atg cag atg acc ctc aag cga tgc    293
73         Glu Cys Ser Val Cys Lys Tyr Lys Met Gln Met Thr Leu Lys Arg Cys
74                  75                  80                  85
75         aag cac ttc gag ctt gga gga gac aag aag acc aag gga gcc gcc atc    341
76         Lys His Phe Glu Leu Gly Gly Asp Lys Lys Thr Lys Gly Ala Ala Ile
77              90                  95                  100
78         tct ttc taa                                                       350
79         Ser Phe
80         105
82 <210> SEQ ID NO: 3
83 <211> LENGTH: 106
84 <212> TYPE: PRT
85 <213> ORGANISM: Phaffia rhodozyma
86 <400> SEQUENCE: 3
87         Met Val Asn Val Pro Lys Thr Arg Arg Thr Tyr Cys Lys Gly Lys Ala
88          1               5                  10                  15
89         Cys Lys Lys His Thr Pro His Lys Val Thr Gln Tyr Lys Lys Gly Lys
90                      20                  25                  30
91         Asp Ser Ile Phe Ala Gln Gly Lys Arg Arg Tyr Asp Arg Lys Gln Ser
92                  35                  40                  45
93         Gly Tyr Gly Gly Gln Thr Lys Pro Val Phe His Lys Lys Ala Lys Thr
94              50                  55                  60
95         Thr Lys Lys Val Val Leu Arg Leu Glu Cys Ser Val Cys Lys Tyr Lys
```

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 96        65                  70                  75                  80
 97        Met Gln Met Thr Leu Lys Arg Cys Lys His Phe Glu Leu Gly Gly Asp
 98                            85                  90                  95
 99        Lys Lys Thr Lys Gly Ala Ala Ile Ser Phe
100                  100                 105
102 <210> SEQ ID NO: 4
103 <211> LENGTH: 741
104 <212> TYPE: DNA
105 <213> ORGANISM: Phaffia rhodozyma
106 <220> FEATURE:
107 <221> NAME/KEY: misc_feature
108 <222> LOCATION: (0)...(0)
109 <223> OTHER INFORMATION: n=a, t, c, or g
110 <400> SEQUENCE: 4
111        ctcgagtgga cggtggcaat ggcattcgtg tcgttggtgc tcactcgcaa cccaagcagt 60
112        cgcttacccg gggtagcctc cgggtgggcg cgatgatttg tggtgtggat tccttcccta 120
113        tgggtagaac gacgcgcaac caatcattcg gagaaccgct ccgttgtagc cgaccagtct 180
114        gattgatcaa catgccagca cgtcctccgg gacggagact ggcggggatc gtacctcatc 240
115        tggaatcgct ggctcaatgg tagtagtctt cacgatcggc catgagggca gtctaggtgg 300
W--> 116   gttcgcctgc cgaagactgt gtgagtgtgc tganaactaa ttgagtaccg ggggataagg 360
117        caaggcgtgt ntggttgccg gtggctgtga gcgagtttgc tgcaaagcga ttcaatgcac 420
118        cccggcttgg ccagcgcgct gcgtcacgaa acacactaaa cggttgacgc cataaagtaa 480
119        taacacactc aagtttgtgg tcccgggtgg gcctctgtgc ctgcgtggga cccgacggga 540
120        gaggaaaacg ttctgtggcc ctctcctctg tggatagtta cctggttgat cctgccagta 600
121        gtcatatgct tgtctcaaag attaagccat gcatgtctaa gtataaacaa attcatactg 660
122        tgaaactgcg aatggctcat taaatcagtt atagtttatt tgatggtacc ttgctacatg 720
123        gataactgtg gtaattctag a                                         741
125 <210> SEQ ID NO: 5
126 <211> LENGTH: 23
127 <212> TYPE: DNA
128 <213> ORGANISM: Artificial Sequence
129 <220> FEATURE:
130 <223> OTHER INFORMATION: CYH1, a PCR primer for the cloning of L41 genomic
131        DNA fragment
132 <220> FEATURE:
133 <221> NAME/KEY: misc_feature
134 <222> LOCATION: (0)...(0)
135 <223> OTHER INFORMATION: n=a, t, c, or g
136 <400> SEQUENCE: 5
W--> 137   cgcgtagtta aygtnccnaa rac                                        23
139 <210> SEQ ID NO: 6
140 <211> LENGTH: 25
141 <212> TYPE: DNA
142 <213> ORGANISM: Artificial Sequence
143 <220> FEATURE:
144 <223> OTHER INFORMATION: CYH3, a PCR primer for the cloning of L41 genomic
145        DNA fragment
146 <400> SEQUENCE: 6
147        cccgggtytt ggcyttyttr tgraa                                     25
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RAW SEQUENCE LISTING   DATE: 09/08/2003
PATENT APPLICATION: **US/09/830,691A**   TIME: 11:27:28

Input Set : **N:\EBONY'S\US09830691A.raw.txt**
Output Set: **N:\CRF4\09082003\I830691A.raw**

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149 <210> SEQ ID NO: 7
150 <211> LENGTH: 24
151 <212> TYPE: DNA
152 <213> ORGANISM: Artificial Sequence
153 <220> FEATURE:
154 <223> OTHER INFORMATION: 3' RACE primer
155 <400> SEQUENCE: 7
156      ggtcagacca agcaagtttt tcac                                    24
158 <210> SEQ ID NO: 8
159 <211> LENGTH: 24
160 <212> TYPE: DNA
161 <213> ORGANISM: Artificial Sequence
162 <220> FEATURE:
163 <223> OTHER INFORMATION: 5' RACE primer
164 <400> SEQUENCE: 8
165      gtgaaaaact tgcttggtct gacc                                    24
167 <210> SEQ ID NO: 9
168 <211> LENGTH: 24
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
171 <220> FEATURE:
172 <223> OTHER INFORMATION: sense primer for the mutagenesis of L41 gene
173 <400> SEQUENCE: 9
174      ggtcagacca agcaagtttt tcac                                    24
176 <210> SEQ ID NO: 10
177 <211> LENGTH: 24
178 <212> TYPE: DNA
179 <213> ORGANISM: Artificial Sequence
180 <220> FEATURE:
181 <223> OTHER INFORMATION: antisense primer for the mutagenesis of L41 gene
182 <400> SEQUENCE: 10
183      gtgaaaaact tgcttggtct gacc                                    24
185 <210> SEQ ID NO: 11
186 <211> LENGTH: 20
187 <212> TYPE: DNA
188 <213> ORGANISM: Artificial Sequence
189 <220> FEATURE:
190 <223> OTHER INFORMATION: a PCR primer corresponding to 18S rDNA
191 <400> SEQUENCE: 11
192      tcctagtaag cgcaagtcat                                         20
194 <210> SEQ ID NO: 12
195 <211> LENGTH: 20
196 <212> TYPE: DNA
197 <213> ORGANISM: Artificial Sequence
198 <220> FEATURE:
199 <223> OTHER INFORMATION: a PCR primer corresponding to 18S rDNA
200 <400> SEQUENCE: 12
201      ttcggccaag gaaagaaact                                         20
203 <210> SEQ ID NO: 13
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RAW SEQUENCE LISTING   DATE: 09/08/2003
PATENT APPLICATION: **US/09/830,691A**   TIME: 11:27:28

Input Set : **N:\EBONY'S\US09830691A.raw.txt**
Output Set: **N:\CRF4\09082003\I830691A.raw**

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204 <211> LENGTH: 20
205 <212> TYPE: DNA
206 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: a PCR primer corresponding to 28S rDNA
209 <400> SEQUENCE: 13
210      aatcggatta tccggagcta                                          20
212 <210> SEQ ID NO: 14
213 <211> LENGTH: 20
214 <212> TYPE: DNA
215 <213> ORGANISM: Artificial Sequence
216 <220> FEATURE:
217 <223> OTHER INFORMATION: a PCR primer corresponding to 28S rDNA
218 <400> SEQUENCE: 14
219      gctataacac atccggagat                                          20
221 <210> SEQ ID NO: 15
222 <211> LENGTH: 2192
223 <212> TYPE: DNA
224 <213> ORGANISM: Phaffia rhodozyma
225 <400> SEQUENCE: 15
226      aagagctatt tgaatgacga ccacaagagt gacgatcata ttgagcatag tataccaaag 60
227      gccaagaggc tgtgtggtgt tctatgagtg gccttgatta tgtgttacat aaataaactg 120
228      atctcaattt ttcaaatact tgccaacact ttcatatatt cacaccaaaa aaagtcagat 180
229      tggcccacaa agtcagatac acgctcgatc gtcgacgggt tcaagcactt tgtcaggcga 240
230      aagaaaggcc acagcaccac ccttcaagtc tcgtctcaat caggttcgtc tagctttttg 300
231      tgtgcaagga tttaccgtct tgatggattt gttcgttgaa agagaggaaa gaacatgctg 360
232      aactgacgaa agtgtgaaca aaaaattgtg attttttcat tgtgtttcgc tggtctcctt 420
233      gctgggttgg gttggatcgg atttatcttc tgtgttggat ggaaaaccct gaatgttctt 480
234      ttcttggaca tcttctaaac tcgacaaaac gattcattcc tccgtactgc tctggttctg 540
235      cctttttgaa tcgcatcgat aaattcttcc ctcggaacgt tcgatcaatc tccgtcaaac 600
236      ttatcatcca aaaatctctt ctcgactgcc gccttgctcc ttttcttcgt tctttcctta 660
237      atccgctttc gactaccctc cttctcttca cactcatagt caagatggtc aacgttccca 720
238      agactcgacg tgagttatag caatttcaac aactctccag acgacaaata ttccagtgca 780
239      tcgaaagagt ttgtggataa acgcgacagt ttcaagggaa agagtcgatg gacagatttg 840
240      gaagacttag ccggtcaagg aacttgggga tcacgtggcg gaggactcat cagaagaagt 900
241      cgggatttgt ttgatcatag tgggatcaag acaaactgga ggatatggct cgccttggaa 960
242      gggaatctcc ggcctggatt cgaggatccg aaagttgtac gtatggaaaa gcttacacgg 1020
243      cttggattta ttatctttca taggaaccta ctgcaagggt aaggcttgca agaagcacac 1080
244      gtaagtcgct tatcctctcc actctttcat ggcatattgt caacgactgg acaacgcgtc 1140
245      cgttttgaaa caagtgactt acctgtgaaa tttgattcta cacctgtatt tagccctcac 1200
246      aaggtacata tcacatcctc ccaccccacc ctgcccaact tcttcagttc atcttgctct 1260
247      cggtttccac attccctgat gacctccttg tatgttcttt gcgaacgttt gtttctgttt 1320
248      ctgtaggtga cccagtacaa gaagggaaag gactccatct tcgcccaggg aaagcgacga 1380
249      tacgaccgaa agcagtccgg ttacgaggt cagaccaagc ccgtttttca caagaaggct 1440
250      aagaccacca agaaggtcgt ccttcgattg ggtacgtttt tgtttatttt gaattctttt 1500
251      tgtgtatgca gacttttgat gattatgctc ctctgtcgtt ttttctcttc aaacagagtg 1560
252      ctccgtctgc agttcgttct tccttccaac caaaacttca actacagaca tcataaacag 1620
253      acatcttact tcggtgttct ctcttttttt ccgcagagta caagatgcag atgaccctca 1680
254      agcgatgcaa gcacttcgag cttggaggag acaagaagac caagggttcg tcttttgtcc 1740
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 09/08/2003
PATENT APPLICATION: **US/09/830,691A**   TIME: 11:27:29

Input Set : **N:\EBONY'S\US09830691A.raw.txt**
Output Set: **N:\CRF4\09082003\I830691A.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; N Pos. 334,371
Seq#:5; N Pos. 15,18

VERIFICATION SUMMARY   DATE: 09/08/2003
PATENT APPLICATION: **US/09/830,691A**   TIME: 11:27:29

Input Set : **N:\EBONY'S\US09830691A.raw.txt**
Output Set: **N:\CRF4\09082003\I830691A.raw**

L:13 M:270 C: Current Application Number differs, Wrong Format
L:116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:300
M:341 Repeated in SeqNo=4
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0

STATISTICS SUMMARY   DATE: 09/08/2003
PATENT APPLICATION: **US/09/830,691A**   TIME: 11:27:29

Input Set : **N:\EBONY'S\US09830691A.raw.txt**
Output Set: **N:\CRF4\09082003\I830691A.raw**

Application Serial Number: US/09/830,691A
Alpha or Numeric or Xml: Numeric
Application Class:
Application File Date: 04-26-2001
Art Unit: PCT
Software Application: FastSEQ
Total Number of Sequences: 20
Total Nucleotides: 4766
Total Amino Acids: 224
Number of Errors: 0
Number of Warnings: 3
Number of Corrections: 1

MESSAGE SUMMARY

270 C: 1 (Current Application Number differs)
341 W: 3 ((46) "n" or "Xaa" used)